

Figure 2

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G(olf) Galphas XLG(olf) XLGalphas	
G(olf) Galphas XLG(olf) XLGalphas	MGCLGGNSKTTEDQGVDEKEMGCLGGNSKTTEDQGVDEKE TARTLLPRGGEGSPACARPKADKPKEKRQRTEQLSAEEREAAKEREAVKE KPQRNLLRNFLVQAFGGCFGRSESPQPKASRSLKVKKVPLAEKRRQMRKEALEKRAQKRA
G(olf) Galphas XLG(olf) XLGalphas	βγ subunit binding RREANKKIEKQLQKERLAYKATHRLLLLGAGESGKSTIVKQMRILHVNGFNPEE QREANKKIEKQLQKDKQVYRATHRLLLLGAGESGKSTIVKQMRILHVNGFNGEGEEDPQ ARKVSRGIDRMLRDQKRDLQQTHRLLLLGAGESGKSTIVKQMRILHVNGFNPEE EKKRSKLIDKQLQDEKMGYMCTHRLLLLGAGESGKSTIVKQMRILHVNGFNGEGGEEDPQ EKKRSKLIDKQLQDEKMGYMCTHRLLLLGAGESGKSTIVKQMRILHVNGFNGEGGEEDPQ
G(olf) Galphas XLG(olf) XLGalphas	KKQKILDIRKNVKDAIVTIVSAMSTIIPPVPLANPENQFRSDYIKSIAPIT AARSNSDGEKATKVQDIKNNLKEAIETIVAAMSNLVPPVELANPENQFRVDYILSVMNVP KRQKILDIRKNVKDAIVTIVSAMSTIIPPVPLANPENQFRSDYIKSIAPIT AARSNSDGEKATKVQDIKNNLKEAIETIVAAMSNLVPPVELANPENQFRVDYILSVMNVP

Figure 4

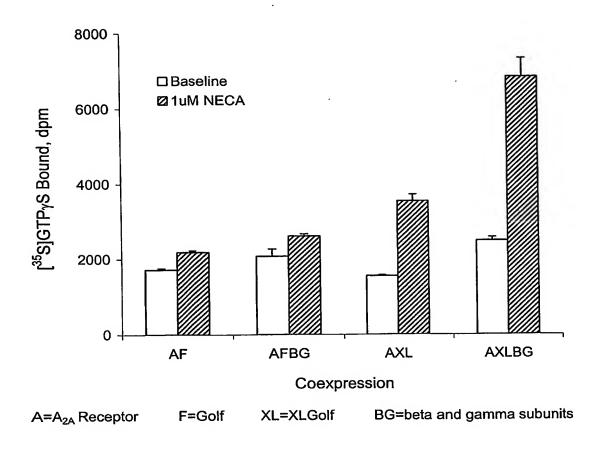


Figure 5

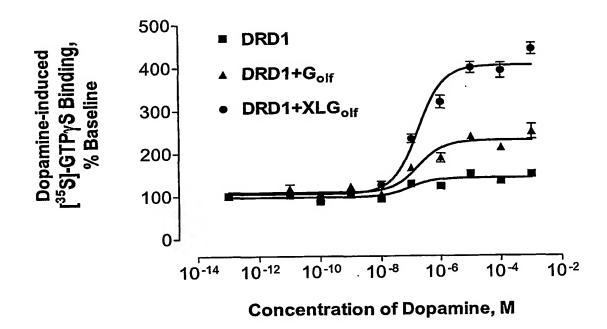


Figure 6

T98G Glioma DNA Golf and XL Golf MSP

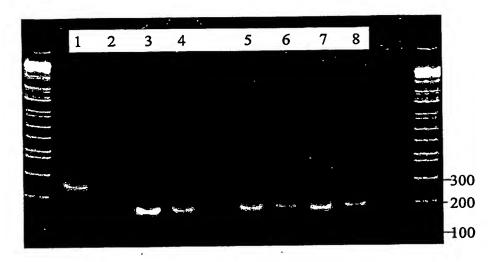
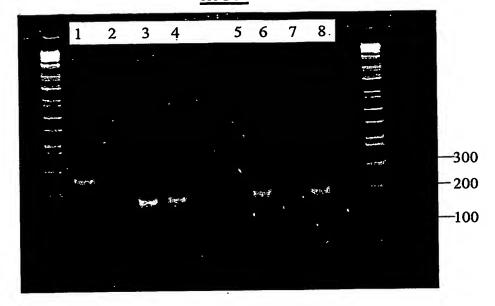


Figure 7

Peripheral Blood DNA Golf and XL Golf MSP



WO 2005/047318 PCT/GB2004/004749

Figure 8

Human GNAL locus; 197140 nt

Features	
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CpG island	21182881
Variation T/C	2198
Variation A/G	2216
Variation C/T	2268
XLGolf start codon	2309
Variation repeat	24212426
Variation A/G	2445
Variation repeat	27462765
Variation C/T	2765
CpG island	6425365647
Variation T/G	64734
Variation C/T	64748
Exon 1	6492565323
Golf start codon	65179
Exon 2	6559865670
Exon 3	6637366427
Exon 4	6657166690
Exon 5	137663137760
Exon 6	175140175194
Exon 7	177278177351
Exon 8	179913179971
Exon 9	181288181408
Exon 10	185013185143
Exon 11	189366189433
Exon 12	193734194571
Stop codon	193878
-	

gagcccagga gttctatatt gcagtgagct atgatcatgc cactgcattc cagcctgggt 60 gacagagaga gatcctgtct taaaaaaaaa aatccataaa atatttgttt tcattttcag 120 ctgactttag tatacaaaat attatccttc ttgtatgata ttgataaagt ttagctaata 180 agttataata gaaaagcaat tgctgtaagt ctcttgggtc tcagtttctt caaaagcctt 240 tectgeetea cacactaete ttecceatet egatttaaac aggacettee tttataetet 300 gagaatcetg tttttccct tcatggcatt agcgtaattt acaattaata tatttacctg 360 tgtttttatg attgagtata tttctacctc attagtctat aaattccatg aggacccctg 420 tetttgeete acceageace aacaagaatg cetagggtac tgtaggcact taattaaatg 480 540 tagtaacttt ttctattttc ccaagttaga ttttctatag tcctcctttc ttttgctcaa 600 atatctaaaa gtatgccata attttagcaa aatttgggga acaatgttag gtcaaaagta 660 gtacatgtat gcacatttgt tatcagtagg acccccaaaa gaaatgtgaa tgccggatct 720 ccaacttett gatttaaaaa tgtaatccag gcccggcgcg ctggctcacg cctgtaatcc 780 cagcactttg ggaggcaagg tgggtggatc cctcgaggtc aggagttcga gaccagcagc 840 ctggccaaca tggtgaaacc gtctctacta aaaattcaaa aaaaaaaaga aaaaaagaaa 900 aattagetgg gagtggtggt gggtgeetge agteceaget actegggagg etgaggeagt 960 agaaccactg gaaccgggga ggcggaggtt gcagtgagcc gtgattgcac cactgcactc 1020 tagcctgggc aacagagcca gactgtttca aaaagaaaaa atgtaatcca atgtagtatt 1080 tacatctagt gccaacgggt acagtgcaca ctgtgctgca tgctgtgttt cattaagcag 1140 cctaatctgg tgtgttaatg agagatgtat ttaactatca gctataatat tccatgtcat 1200 tttccaagga acacatctta cagcaggttt ttcacaagct atattgaaat gttcacctgt 1260 tgcagaagct ctataagatg ctatgcattc agcgggaaat aaccgaagac atagcacctc 1320 tggcaggett gtttctcage gtcatggaga gagegcacce agtctgaagg egeatctget 1380 cetetetgee ctatattgtg gattaagaaa atacagtgta taateteata tteteattte 1440 agcaaatata aatagtacat ggcaattata tgtgctcggt tttcattttt aagggtggag 1500 attgttgaaa atggtgtcgt ggaccagact ccagaattgg agattttgta gagatcaaag 1560

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#### Fig. 8 (cont.)

Fig. 8 (cont.)						
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#### Fig. 8 (cont.)

g. 8 (cont.)						
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Fig. 8 (cont.)						
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## Fig. 8 (cont.)

Fig. 8 (cont.)						
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tttctctgac	ctatccqtqc	tacttcaacc	tcctgggggc	aatgataagg	gtgaggttat	193560
ctagatecte	aaccaatact	tgcattgaga	ccattcctgc	ctctaagtgc	tcccatgcaa	193620
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ccagagtaca	tactagaacc	gcgcagggct	agtgcacacg	ctctctcttg	cagaggatca	193740
gcacggccac	caataacaac	aaacattact	gctacccgca	cttcacctgc	geegtggaea	193800
cagagaacat	ccacaaaata	ttcaacgact	gccgcgacat	catccagcgg	atgeacetea	193860
agcagtatga	actettataa	ggatgctgcc	gccaccctgc	gacggagcgg	cgccccggac	193920
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tecetttatt	gattcatcga	ggagaacttg	gtagatgggg	agaaaacaca	gilggilli	194220
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acattcagag	tegetgetgg	cctcaggcat	ttgaattaga	gctactttga	gcctcttagg	194520
cagaaaacct	accacattca	ctactgcaaa	atgtgtcctg	tctaaaaatg	attctctaaa	194580
ctttccctat	. acttaggcat	agtcttcttt	cttagattct	ctttgttgtt	gtccctattg	194640
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ctgatttaaa	. aaattgtaag	ttatacacgt	taatcatcca	cattetateg	acaatgtacc	194940
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taagcaaaca	ı gtaagagaaa	aacaaaatgt	. ggccaggcgc	. gg.ggc.cac	gcctgtaatc	195240
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ggcacctgta	atctcagcta	cucyayaygc	. Lyayycayyd	. gaalceceg	aacctgggag tcaaagtgag	195420
gtggagattg	g cagtgagccg	aggicgige	. accycacicc	, agoolggaca · castatttat	gtctggccta	195480
actcaggcca	aaaaaaaaa	taacttacct	tttaaagaa	. caacycecyc	aaaccttaat	195540
ggagaatgag	, gatgacagct	. Coactigeet	. cityaayaay	, addicated		

g. o (conc.)						
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cctcaaacat	tatgacacac	caacaatatc	tagaaagtaa	gaactggtgt	agcaatgatg	195660
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totataaaga	tatccasaat	taaattttca	acaatacaaa	tctagagaag	tgacagccta	195780
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cattatttta	atttataat	cctttataaa	ctcatccaga	tttaaatgct	actttttcat	195900
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aacaaaaaga	acacttccca	tttttaaatt	tottttatt	actgacaggc	attaacagat	196320
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gtagcaacgt	ggteteetat	agagaaaacc	atattataca	catcacatct	gtacttttt	196440
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tttttaaat	atattttttg	agacggagcc	teacceage	aagggattct	cctacctcaa	196560
gcatgatctt	ggeteactge	aacctccgcc	atacttacet	aatttttgta	tttctagcgg	196620
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gatttgttgt	agagtacctg	tccactttta	tagcatgaga	acagtacaat	caactattta	197100
ttttgcagtt	actcatttca	gtgattgaga	atttctgtgc	tgtgcagaga	gacggcctgt	197140
aattggtctc	atcatccact	tgattctaac	atgatctctg			19/140

## Figure 9

## Amino Acid Sequence of Human Golf

_			DDDDNWATER	QLQKERLAYK	ATHIDIT.T.T.CA	CESCKSTIVK
1	MGCLGGNSKT	TEDQGVDEKE	KKEANKKIEK	<b>ÖTÖVEYTY 1 V</b>	WILLUMBERGY	ODDOMBILVI
61	OMPTI.HUNGE	NPEEKKOKTI	DTRKNVKDAT	VTIVSAMSTI	IPPVPLANPE	NQFRSDYIKS
0.7	<b>MALTINAMOT</b>	TAT DELGASTATE	22144111411			CITCI INVIDE
121	TADITOREVS	OEFFDHVKKL	WDDEGVKACF	ERSNEYQLID	CAGALPERID	SASPADILLI
121	THETTOPIN	22112111			****	TTUTTA ACCCU
181	DODLLRCRVL	TSGIFETRFQ	VDKVNFHMFD	VGGQRDERIK	MIQCENDVIA	TTIVAACSSI
-01	DQDEE:::			MTOTTI DI NIK	ODME A PERSON	CKCKIEDAED
241	NMVIREDNNT	NRLRESLDLF	ESIWNNRWLR	TISIILFLNK	<b>ÖDMTYEVADY</b>	GYPKTEDILE
				T DT DT COD COC	DCKRACADRE	TOATHTENTE
301	EYANYTVPED	ATPDAGEDPK	VIRAKFFIRD	LFLRISTATG	DGKHICIFIII	ICHADIMATIC
361	RVFNDCRDII	OKMMPKÖXEP	ш			

## Figure 10

## cDNA and Amino Acid Sequence of Human XLGolf

	MGLCYSLRPLLFGGPGDDPC
1	ATGGGTCTGTGCTACAGTCTGCGGCCGCTGCTTTTCGGGGGGCCCAGGGGACGACCCCTGC
	AASEPPVEDAQPAPALAP
61	GCGGCCTCGGAGCCGCCGGTGGAGGACGCGCAGCCCCGGCCCCGGCCCTGGCCCCA
	V R A A A R D T A R T L L P R G G E G S
121	GTCCGGGCGGCCGCAAGGGACACGGCCCGGACCCTGCTCCCTCGGGGCGGCGAAGGGAGC
	PACARPKADKPKE KRQRTEQ
181	CCGGCATGCGCTCGGCCCAAAGCAGACAAGCCGAAGGAGAAGCGGCAGCGCACCGAGCAG
	LSAEEREAAKEREAVKEARK
241	CTGAGTGCCGAGGAGCGCGAGGCGCCAAGGAGCGCGAGGCGAGGAAA
	V S R G I D R M L R D Q K R D L Q Q T H
301	GTGAGCCGGGGCATCGACCGCATGCTGCGCGACCAGAAGCGCGACCTGCAGCAGACGCAC
	R·L L L G A G E S G K S T I V K Q M R
361	CGGCTCCTGCTCGGGGCTGGTGAGTCTGGGAAAAGCACCATCGTCAAACAGATGAGG
	T, D V A M G L M L D D V V & W D D D D D D D D D D D D D D D D D D
421	ATCCTGCACGTCAATGGGTTTAATCCCGAGGAAAAGAAACAGAAAATTCTGGACATCCGG
	K N V K D A I V T I V S A M S T I I P P AAAAATGTTAAAGATGCTATCGTGACAATTGTTTCAGCAATGAGTACTATAATACCTCCA
481	
	V P L A N P E N Q F R S D Y I K S I A P GTTCCGCTGGCCAACCCTGAAAACCAATTTCGATCAGACTACATCAAGAGCATAGCCCCT
541	
C01	I T D F E Y S Q E F F D H V K K L W D D ATCACTGACTTTGAATATTCCCAGGAATTCTTTGACCATGTGAAAAAACTTTGGGACGAT
601	E.G.V.K.A.C.F.E.R.S.N.E.Y.Q.L.I.D.C.A.Q
661	GAAGGCGTGAAGGCATGCTTTGAGAGATCCAACGAATACCAGCTGATTGACTGTGCACAA
991	Y F L E R I D S V S L V D Y T P T D Q D
721	TACTTCCTGGAAAGAATCGACAGCGTCAGCTTGGTTGACTACACACCCACAGACCAGGAC
121	L L R C R V L T S G I F E T R F Q V D K
781	CTCCTCAGATGCAGAGTTCTGACATCTGGGATTTTTGAGACACGATTCCAAGTGGACAAA
701	V N F H M F D V G G Q R D E R R K W I Q
841	GTAAACTTCCACATGTTTGATGTTGGTGGCCAGAGGGATGAGAGGAGAAAATGGATCCAG
	C F N D V T A I I Y V A A C S S Y N M V
901	TGCTTTAACGATGTCACAGCTATCATTTACGTCGCAGCCTGCAGTAGCTACAACATGGTG
	I R. E D N N T N R L R E S L D L F E S I
961	ATTCGAGAAGATAACAACACCAACAGGCTGAGAGAGTCCCTGGATCTTTTTGAAAGCATC
	WNNRWLRTISIILFLNKQDM
1021	TGGAACAACAGGTGGTTACGGACCATTTCTATCATCTTGTTCTTGAACAAACA
	LAEKVLAGKSKIEDYFPEYA
1081	CTGGCAGAAAAGTCTTGGCAGGGAAATCAAAAATTGAAGACTATTTCCCAGAATATGCA
	N Y T V P E D A T P D A G E D P K V T R
1141	AATTATACTGTTCCTGAAGACGCAACACCAGATGCAGGAGAAGATCCCAAAGTTACAAGA
	A K F F I R D L F L R I S T A T G D G K
1201	GCCAAGTTCTTTATCCGGGACCTGTTTTTGAGGATCAGCACGGCCACCGGTGACGGCAAA
1261	CATTACTGCTACCCGCACTTCACCTGCGCCGTGGACACAGAGAACATCCGCAGGGTGTTC
1321	AACGACTGCCGCGACATCATCCAGCGGATGCACCTCAAGCAGTATGAGCTCTTGTGA

PCT/GB2004/004749

## Figure 11

G _{olf} G _{as} XLG _{olf} XLG _{as}	MEISGPPFEIGSAPAGVDDTPVNMDSPPIALDGPPIKVSGAPDKRERAERPPVEEEAAEM
G _{olf} G _{αs} XLG _{olf} XLG _{αs}	EGAADAAEGGKVPSPGYGSPAAGAASADTAARAAPAAPADPDSGATPED PDSGTAPADPD
G _{olf} G _{αs} XLG _{olf} XLG _{αs}	SGAFAADPDSGAAPAAPADPDSGAAPDAPADPDSGAAPDAGAAPEAPAAPAAAE
G _{olf} G _{as} XLG _{olf} XLG _{as}	TRAAHVAPAAPDAGAPTAPAASATRAAQVRRAASAAPASGARRKIHLRP PSPEIQAADPP
G _{olf} G _{as} XLG _{olf} XLG _{as}	TPRPTRASAWRGKSESSRGRRVYYDEGVASSDDDSSGDESDDGTSGCLRWFQHRRNRRR
G _{olf} G _{os} XLG _{olf} XLG _{os}	MGCLGGNSKTTEDQGVDEKEMGCLG-NSKT-EDQRNEEKA LAPVRAAARDTARTLLPRGGEGSPACARPKADKPKEKRQRTEQLSAEER EAAKEREAVKE KPQRNLLRNFLVQAFGGCFGRSESPQPKASRSLKVKKVPLAEKRRQMRK EALEKRAQKRA
G _{olf} G _{αs} XLG _{olf} XLG _{αs}	βγ binding domain RREANKKIEKQLQKERLAYKATHRLLLLGAGESGKSTIVKQMRILHVNGFNPE QREANKKIEKQLQKDKQVYRATHRLLLLGAGESGKSTIVKQMRILHVNGFNGE ARKVSRGIDRMLRDQKRDLQQTHRLLLLGAGESGKSTIVKQMRILHVNGFNPE EKKRSKLIDKQLQDEKMGYMCTHRLLLLGAGESGKSTIVKQMRILHVNGFNGE

## Figure 12

	1			50
G(olf) XL-G(olf)	atgggtctgt gctaca	gtct gcggccgctg		
0/-15)	51			100
G(olf) XL-G(olf)	cgacccctgc gcggcc	tegg ageegeeggt	ggaggacgcg	cagcccgccc
G(olf)	101	,		150
XL-G(olf)	eggeeeegge eetgge	ccca gtccgggcgg	ccgcaaggga	
G(olf)	151	,		200
XL-G(olf)	accetgetee cteggg	ggcgg cgaagggagc	ccggcatgcg	eteggeecaa 250
G(olf) XL-G(olf)	agcagacaag ccgaag	ggaga agcggcagcg	~ <u>AtaGaGtat</u> cAccGaGcag	tTGqGcGqCa cTGaGtGcCg
G(olf) XL-G(olf)	251 <u>Acaqcaaqac GaCGG</u> Aggagcgcga GgCGG	aaqAc cAGqGCGtcG ccaAg gAGcGCGagG	atGaaAAaGA cgGtcAAgGA	300 <u>acqacGcqAq</u> ggcgaGgaAa
G(olf) XL-G(olf)	301 <u>GccAaCaaaa aqATC</u> GtgAgCcggg gcATC	GAqaa qcaGtTGCac GAccg catGcTGCgc	aAaqAGcqcC gAccAGaagC	350 tgGcttacaA gcGacctgcA
G(olf) XL-G(olf)	351 <u>GgctACcCAC CGcCT</u> GcagACgCAC CGgCT	<u>aCTGC TcCTqG</u> GGGG cCTGC TgCTcGGGGG	TGGTGAGTCT TGGTGAGTCT	400 GGGAAAAGCA GGGAAAAGCA
G(olf) XL-G(olf)	401 CTATCGTCAA ACAGA CTATCGTCAA ACAGA	FGAGG ATCCTGCACG	TCAATGGGTT TCAATGGGTT	450 TAATCCCGAG TAATCCCGAG
G(olf) XL-G(olf)	451 GAAAAGAAAC AGAAA GAAAAGAAAC AGAAA	ATTCT GGACATCCGC	AAAAATGTTA AAAAATGTTA	500 AAGATGCTAT AAGATGCTAT
G(olf) XL-G(olf)	501 CGTGACAATT GTTTC CGTGACAATT GTTTC	AGCAA TGAGTACTAT AGCAA TGAGTACTAT	ADDTDDATAA T ADDTDDATAA T	550 GTTCCGCTGG GTTCCGCTGG
G(olf) XL-G(olf)	551 CCAACCCTGA AAACC CCAACCCTGA AAACC	AATTT CGATCAGACT	T ACATCAAGAG T ACATCAAGAG	600 CATAGCCCCT CATAGCCCCT
G(olf) XL-G(olf)	601 ATCACTGACT TTGAA ATCACTGACT TTGAA	TATTC CCAGGAATT( TATTC CCAGGAATT(	C TTTGACCATG	650 TGAAAAACT TGAAAAAACT
G(olf) XL-G(olf)	651 TTGGGACGAT GAAGG TTGGGACGAT GAAGG	CGTGA AGGCATGCT CGTGA AGGCATGCT	r tgagagatcc r tgagagatcc	700 AACGAATACC AACGAATACC
G(olf) XL-G(olf)	701 AGCTGATTGA CTGTG AGCTGATTGA CTGTG	CACAA TACTTCCTGG	AAAGAATCGA AAAGAATCGA	750 CAGCGTCAGC CAGCGTCAGC
G(olf) XL-G(olf)	751 TTGGTTGACT ACACA TTGGTTGACT ACACA	AGDACCAGA OACCO. AGACCAGA AGACCAGO	C CTCCTCAGAT C CTCCTCAGAT	800 GCAGAGTTCT GCAGAGTTCT
G(olf) XL-G(olf)	801 GACATCTGGG ATTTT GACATCTGGG ATTTT	TGAGA CACGATTCC: TGAGA CACGATTCC:	A AGTGGACAAA A AGTGGACAAA	850 GTAAACTTCC GTAAACTTCC
G(olf) XL-G(olf)	851 ACATGTTTGA TGTTG ACATGTTTGA TGTTG	GTGGC CAGAGGGAT GTGGC CAGAGGGAT	G AGAGGAGAAA G AGAGGAGAAA	900 ATGGATCCAG ATGGATCCAG
G(olf) XL-G(olf)	901 TGCTTTAACG ATGTC TGCTTTAACG ATGTC	ACAGC TATCATTTA ACAGC TATCATTTA	C GTCGCAGCCT C GTCGCAGCCT	950 GCAGTAGCTA GCAGTAGCTA
G(olf) XL-G(olf)	951 CAACATGGTG ATTCG CAACATGGTG ATTCG	AGAAG ATAACAACA AGAAG ATAACAACA	C CAACAGGCTG C CAACAGGCTG	1000 AGAGAGTCCC AGAGAGTCCC

G(olf) XL-G(olf)	1001 TGGATCTTTT TGGATCTTTT	TGAAAGCATC TGAAAGCATC	TGGAACAACA TGGAACAACA	GGTGGTTACG GGTGGTTACG	1050 GACCATTTCT GACCATTTCT
G(olf) XL-G(olf)	1051 ATCATCTTGT ATCATCTTGT	TCTTGAACAA TCTTGAACAA		CTGGCAGAAA CTGGCAGAAA	
G(olf) XL-G(olf)		DAADTTGAAG DAADTTAAAA	ACTATTTCCC ACTATTTCCC	AGAATATGCA AGAATATGCA	
G(olf) XL-G(olf)		CGCAACACCA CGCAACACCA		AAGA†CCCAA AAGA¢CCCAA	
G(olf) XL-G(olf)	1201 GCCAAGTTCT GCCAAGTTCT	TTATCCGGGA TTATCCGGGA		AGGATCAGCA AGGATCAGCA	
G(olf) XL-G(olf)		CATTACTGCT CATTACTGCT		CACCTGCGCC CACCTGCGCC	
G(olf) XL-G(olf)		CAGGGTGTTC CAGGGTGTTC		GCGACATCAT GCGACATCAT	
G(olf) XL-G(olf)		AGTATGAGCT AGTATGAGCT		•	

## Figure 13

					E 0
	1				50
G(olf) $xl-G(olf)$	mglcyslrpl	lfggpgddpc	aaseppveda	qpapapalap	vraaardtar
G(olf) xl-G(olf)	51 ~~~~~~ tllprggegs	~~~~~~ pacarpkadk	<u>mqc</u> pkekrqrteq	<u>Lggnsktted</u> Lsaeereaak	100 ggvdeKErRe ereavKEaRk
G(olf) xl-G(olf)	101 <u>ankklekgLg</u> vsrgIdrmLr	kerlaykaTH dqkrdlqqTH	RLLLLGAGES RLLLLGAGES	GKSTIVKQMR GKSTIVKQMR	150 ILHVNGFNPE ILHVNGFNPE
G(olf) xl-G(olf)	151 EKKQKILDIR EKKQKILDIR	KNVKDAIVTI KNVKDAIVTI	VSAMSTIIPP VSAMSTIIPP	VPLANPENQF VPLANPENQF	200 RSDYIKSIAP RSDYIKSIAP
G(olf) xl-G(olf)	201 ITDFEYSQEF ITDFEYSQEF	FDHVKKLWDD FDHVKKLWDD	EGVKACFERS EGVKACFERS	NEYQLIDCAQ NEYQLIDCAQ	250 YFLERIDSVS YFLERIDSVS
G(olf) xl-G(olf)	251 LVDYTPTDQD LVDYTPTDQD	LLRCRVLTSG LLRCRVLTSG		VNFHMFDVGG VNFHMFDVGG	300 QRDERRKWIQ QRDERRKWIQ
G(olf) xl-G(olf)	301 CFNDVTAIIY CFNDVTAIIY	VAACSSYNMV VAACSSYNMV	IREDNNTNRL IREDNNTNRL	RESLDLFESI RESLDLFESI	
G(olf)	351 IILFLNKQDM IILFLNKQDM	LAEKVLAGKS LAEKVLAGKS	KIEDYFPEYA KIEDYFPEYA	NYTVPEDATP NYTVPEDATP	400 DAGEDPKVTR DAGEDPKVTR
G(olf) xl-G(olf)	401 AKFFIRDLFI AKFFIRDLFI	RISTATGDGK RISTATGDGK	НҮСҮРНГТСА НҮСҮРНГТСА	VDTENIRRVF VDTENIRRVF	450 NDCRDIIQRM NDCRDIIQRM
G(olf) xl-G(olf)	451 HLKQYELL HLKQYELL	•			

## Figure 14

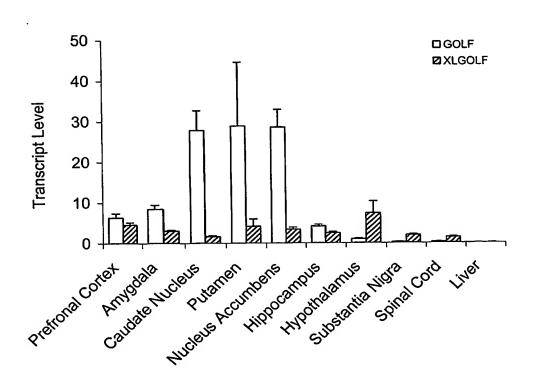


Figure 15

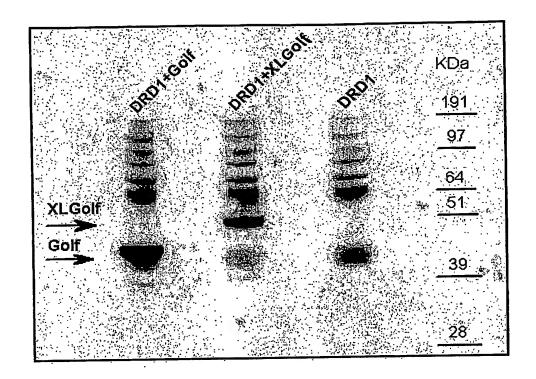
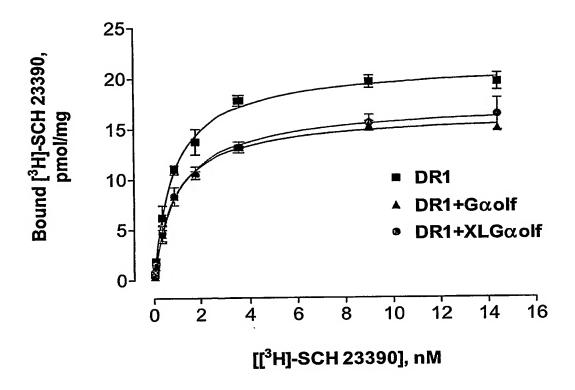
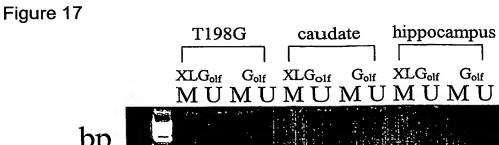
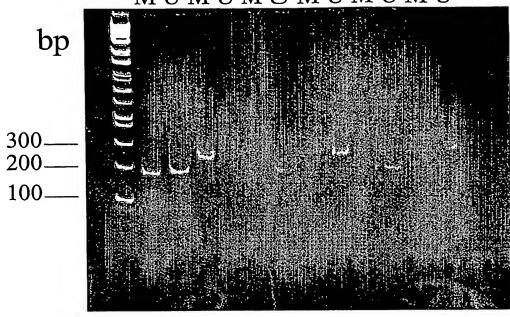
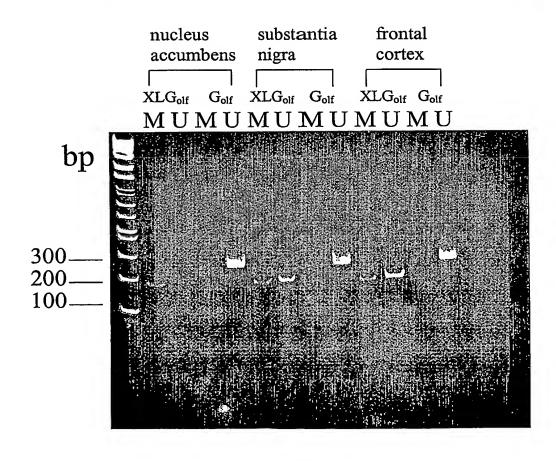


Figure 16









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SUBSTITUTE SHEET (RULE 26)

## Figure 18 Mouse XLGolf cDNA

ggcgaggctcgcgcctgggacgttcgaaggatgcacaatggcccttcggggcagtgagggc cgtggtccccagccggacccaccgcagccacgcccgcagctggcgccccagggtccgagt gcagcgctgcctccggcgtcggcggagccccggtctgacccgcggggacttgcccgcgccc cgcgccccgatgggcctatgctacagcctgcggccgctgctcttcgggagcccagaggacac cccgtgtgcggcctcggaaccctgcgcagaggatgctcagcccagcgccgccccggcccctg cctcgatcccagccccggctcccgtagggaccctgctccggcgtggcggccggatcgtc gcgaacgcgcggccgccaggcgagctgcagagccgccggcgacaggagcagctacgagccga ggagcgcgaggctaaagaggcgaggaaagtcagccggggcatcgaccgcatgctgcgcg agcagaagcgggacctgcagcagacgcaccggctcctgctgctgggggctggtgagtccggg aaaagcactatcgtcaaacagatgaggatcctgcacgtcaatggcttcaaccccgaggaaaa gaagcagaaaattctggacatcaggaaaaatgtcaaagatgcgatcgtgacaatcgtttcag caatgagtactatcatacctccagttccactggccaaccctgagaaccagttccggtcagat tatatcaagagcatagcccctatcactgactttgaatattcccaggagttctttgaccatgt gaagaagctgtgggacgatgaaggagtgaaggcctgctttgagagatccaacgagtaccagc tgatcgactgtgcacaatagtaagttgcttcccaggccaggtctctctgaagcctgattgca ttcttggctgtgcccaccctatagctcagaggtttcctaaaagcagaaatcaagagtaactt tgacttcagaagttaactttctaaagaggaggttttggcaacattgtttccctgaaggtgta ttcagtaatgttgtttcaccaaaggagctggttttggtaacattgtttcactgaaggagatg tgttcagtaacattgtttcacgagcgaaccagagtaagtttctcttttgtcctcttaagttct tttcatttttttttcctatctgcacagagttaaagacatatctactaattataatgtacccaa atattcactattatattttaatttttaagcaagcctctccaaattaagttttttaatatcaa aatgtgcaggactggggagatggcttgctatctataaaagcactggatgcttttccagagga cctgagttcagttcccagcactcagctcacaaccacctataactccagctccaggggatctg gtgccctctgctgacctccttggcacacacaactaattaaaattttaaaaagtctgtttcct attgtttccctgctgcattttggagttcagcagaatgaagcttacattggtctcgggcaggt tcttactccttcgctgctgttttggagtggttccagatttgaaccagcagtgacttcagtgt ccacatactcctggtcatacaactgccctgtaaactagggtacaaagcttcaacagcaactt ttgttatcttgctgtttcctgtgggatgaatttgtgttccaggagaatgatgctcttatatt gtctgagatcaagtacaggccagtgttggatcccacccatagaatccaggtaatttgctttg tgaagaggtttaacctgtcttctctggagctttttgggataaatgagtgtggtgtgagttcc ttccgtgatggttctcaagtaataggacaacattggttgattcccttgcaaagtaaaaactg caaattaacaatccctgtgttaagacccccttcatcccttgagatgcagaaacaagcgaaac ttgctagcctggggcccagtttcaactatactgctattcataccacgaccaatgataacatc agttacctgtttaaatgccttctggggtttggtagaacataactctatagtgtcatcattta atgagttaattctaagtgcactggaactttctctgtgaaggtgaaactcacaataaagctgt tgttgtataagaagaataaataattatttcagg

#### Figure 19

#### Mouse XLGolf Protein

MGLCYSLRPLLFGSPEDTPCAASEPCAEDAQPSAAPAPASIPAPAPVGTLLRRGGGRIVANARPPGELQS' RRRQEQLRAEEREAAKEARKVSRGIDRMLREQKRDLQQTHRLLLLGAGESGKSTIVKQMRILHVNGFN PEEKKQKILDIRKNVKDAIVTIVSAMSTIIPPVPLANPENQFRSDYIKSIAPITDFEYSQEFFDHVKKLWDD EGVKACFERSNEYQLIDCAQ